

SEQUENCE LISTING

- <110> Klee, Harry J. Lashbrook, Coralie Shrode, Lori
- <120> Materials and Methods for Tissue-Specific Targeting of Ethylene Insensitivity in Transgenic Plants
- <130> UF-325XC1
- <140> US 10/602,475
- <141> 2003-06-23
- <150> US 60/390,385
- <151> 2002-06-21
- <160> : 16
- <170> PatentIn version 3.2
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- Phe Pro Tyr Arg Trp Val Leu Val Gln Phe Gly Ala Phe Ile Val Leu 50 55 60
- Tyr Gly Ala Thr His Leu Ile Asn Leu Trp Thr Phe Thr Thr His Ser 65 70 75 80
- Arg Thr Val Ala Leu Val Met Thr Thr Ala Lys Val Leu Thr Ala Val 85 90 95
- Val Ser Cys Ala Thr Ala Leu Met Leu Val His Ile Ile Pro Asp Leu
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- Leu Ser Val Lys Thr Arg Glu Leu Phe Leu Lys Asn Lys Ala Ala Glu 115 120 125
- Leu Asp Arg Glu Met Gly Leu Ile Arg Thr Gln Glu Glu Thr Gly Arg 130 135 140

His Val Arg Met Leu Thr His Glu Ile Arg Ser Thr Leu Asp Arg His 145 150 155 160

2

- Thr Ile Leu Lys Thr Thr Leu Val Glu Leu Gly Arg Thr Leu Ala Leu 165 170 175
- Glu Glu Cys Ala Leu Trp Met Pro Thr Arg Thr Gly Leu Glu Leu Gln 180 185 190
- Leu Ser Tyr Thr Leu Arg His Gln His Pro Val Glu Tyr Thr Val Pro
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- Ile Gln Leu Pro Val Ile Asn Gln Val Phe Gly Thr Ser Arg Ala Val 210 215 220
- Lys Ile Ser Pro Asn Ser Pro Val Ala Arg Leu Arg Pro Val Ser Gly
 225 230 235 240
- Lys Tyr Met Leu Gly Glu Val Val Ala Val Arg Val Pro Leu Leu His
 245 250 255
- Leu Ser Asn Phe Gln Ile Asn Asp Trp Pro Glu Leu Ser Thr Lys Arg 260 265 270
- Tyr Ala Leu Met Val Leu Met Leu Pro Ser Asp Ser Ala Arg Gln Trp 275 280 285
- His Val His Glu Leu Glu Leu Val Glu Val Val Ala Asp Gln Val Ala 290 295 300
- Val Ala Leu Ser His Ala Ala Ile Leu Glu Glu Ser Met Arg Ala Arg 305 310 315 320
- Asp Leu Leu Met Glu Gln Asn Val Ala Leu Asp Leu Ala Arg Arg Glu 325 330 335
- Ala Glu Thr Ala Ile Arg Ala Arg Asn Asp Phe Leu Ala Val Met Asn 340 345 350
- His Glu Met Arg Thr Pro Met His Ala Ile Ile Ala Leu Ser Ser Leu 355 360 365
- Leu Gln Glu Thr Glu Leu Thr Pro Glu Gln Arg Leu Met Val Glu Thr 370 380
- Ile Leu Lys Ser Ser Asn Leu Leu Ala Thr Leu Met Asn Asp Val Leu 385 390 395 400
- Asp Leu Ser Arg Leu Glu Asp Gly Ser Leu Gln Leu Glu Leu Gly Thr 405 410 415
- Phe Asn Leu His Thr Leu Phe Arg Glu Val Leu Asn Leu Ile Lys Pro 420 425 430
- Ile Ala Val Val Lys Lys Leu Pro Ile Thr Leu Asn Leu Ala Pro Asp

435 440 445

Leu Pro Glu Phe Val Val Gly Asp Glu Lys Arg Leu Met Gln Ile Ile 450 455 460

Leu Asn Ile Val Gly Asn Ala Val Lys Phe Ser Lys Gln Gly Ser Ile 465 470 475 480

Ser Val Thr Ala Leu Val Thr Lys Ser Asp Thr Arg Ala Ala Asp Phe 485 490 495

Phe Val Val Pro Thr Gly Ser His Phe Tyr Leu Arg Val Lys 500 505 510

Asp Ser Gly Ala Gly Ile Asn Pro Gln Asp Ile Pro Lys Ile Phe Thr 515 520 525

Lys Phe Ala Gln Thr Gln Ser Leu Ala Thr Arg Ser Ser Gly Gly Ser 530 540

Gly Leu Gly Leu Ala Ile Ser Lys Arg Phe Val Asn Leu Met Glu Gly 545 550 555 560

Asn Ile Trp Ile Glu Ser Asp Gly Leu Gly Lys Gly Cys Thr Ala Ile 565 570 575

Phe Asp Val Lys Leu Gly Ile Ser Glu Arg Ser Asn Glu Ser Lys Gln 580 585 590

Ser Gly Ile Pro Lys Val Pro Ala Ile Pro Arg His Ser Asn Phe Thr 595 600 605

Gly Leu Lys Val Leu Val Met Asp Glu Asn Gly Val Ser Arg Met Val 610 620

Thr Lys Gly Leu Leu Val His Leu Gly Cys Glu Val Thr Thr Val Ser 625 630 635 640

Ser Asn Glu Glu Cys Leu Arg Val Val Ser His Glu His Lys Val Val 645 650 655

Phe Met Asp Val Cys Met Pro Gly Val Glu Asn Tyr Gln Ile Ala Leu 660 665 670

Arg Ile His Glu Lys Phe Thr Lys Gln Arg His Gln Arg Pro Leu Leu 675 680 685

Val Ala Leu Ser Gly Asn Thr Asp Lys Ser Thr Lys Glu Lys Cys Met 690 695 700

Ser Phe Gly Leu Asp Gly Val Leu Leu Lys Pro Val Ser Leu Asp Asn 705 710 715 720

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Phe Pro Tyr Arg Trp Val Leu Val Gln Phe Gly Ala Phe Ile Val Leu 50 55 60

Tyr Gly Ala Thr His Leu Ile Asn Leu Trp Thr Phe Thr Thr His Ser 65 70 75 80

Arg Thr Val Ala Leu Val Met Thr Thr Ala Lys Val Leu Thr Ala Val
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Val Ser Cys Ala Thr Thr Leu Met Leu Val His Ile Ile Pro Asp Leu
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Leu Ser Val Lys Thr Arg Glu Leu Phe Leu Lys Asn Lys Ala Ala Glu
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His Val Arg Met Leu Thr His Glu Ile Arg Ser Thr Leu Asp Arg His 145 150 155 160

Thr Ile Leu Lys Thr Thr Leu Val Glu Leu Gly Arg Thr Leu Ala Leu 165 170 175

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Leu Ser Tyr Thr Leu Arg His Gln His Pro Val Glu Tyr Thr Val Pro 195 200 205

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- Tyr Ala Leu Met Val Leu Met Leu Pro Ser Asp Ser Ala Arg Gln Trp 275 280 285
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- Val Ala Leu Ser His Ala Ala Ile Leu Glu Glu Ser Met Arg Ala Arg 305 310 315 320
- Asp Leu Leu Met Glu Gln Asn Val Ala Leu Asp Leu Ala Arg Arg Glu 325 330 335
- Ala Glu Thr Ala Ile Arg Ala Arg Asn Asp Phe Leu Ala Val Met Asn 340 345 350
- His Glu Met Arg Thr Pro Met His Ala Ile Ile Ala Leu Ser Ser Leu 355 360 365
- Leu Gln Glu Thr Glu Leu Thr Pro Glu Gln Arg Leu Met Val Glu Thr 370 375 380
- Ile Leu Lys Ser Ser Asn Leu Leu Ala Thr Leu Met Asn Asp Val Leu 385 390 395 400
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- Phe Val Val Pro Thr Gly Ser His Phe Tyr Leu Arg Val Lys 500 505 510
- Asp Ser Gly Ala Gly Ile Asn Pro Gln Asp Ile Pro Lys Ile Phe Thr 515 520 525

Lys Phe Ala Gln Thr Gln Ser Leu Ala Thr Arg Ser Ser Gly Gly Ser 530 540

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Asn Ile Trp Ile Glu Ser Asp Gly Leu Gly Lys Gly Cys Thr Ala Ile 565 570 575

Phe Asp Val Lys Leu Gly Ile Ser Glu Arg Ser Asn Glu Ser Lys Gln 580 585 590

Ser Gly Ile Pro Lys Val Pro Ala Ile Pro Arg His Ser Asn Phe Thr 595 600 605

Gly Leu Lys Val Leu Val Met Asp Glu Asn Gly Val Ser Arg Met Val 610 620

Thr Lys Gly Leu Leu Val His Leu Gly Cys Glu Val Thr Thr Val Ser 625 630 635 640

Ser Asn Glu Glu Cys Leu Arg Val Val Ser His Glu His Lys Val Val 645 650 655

Phe Met Asp Val Cys Met Pro Gly Val Glu Asn Tyr Gln Ile Ala Leu 660 665 670

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Val Ala Leu Ser Gly Asn Thr Asp Lys Ser Thr Lys Glu Lys Cys Met 690 695 700

Ser Phe Gly Leu Asp Gly Val Leu Leu Lys Pro Val Ser Leu Asp Asn 705 710 715 720

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- Arg Thr Val Ala Leu Val Met Thr Thr Ala Lys Val Leu Thr Ala Val
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- Leu Ser Val Lys Thr Arg Glu Leu Phe Leu Lys Asn Lys Ala Ala Glu 115 120 125
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- His Val Arg Met Leu Thr His Glu Ile Arg Ser Thr Leu Asp Arg His 145 150 155 160
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- Lys Tyr Met Leu Gly Glu Val Val Ala Val Arg Val Pro Leu Leu His
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- Tyr Ala Leu Met Val Leu Met Leu Pro Ser Asp Ser Ala Arg Gln Trp 275 280 285
- His Val His Glu Leu Glu Leu Val Glu Val Val Ala Asp Gln Val Ala 290 295 300
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8

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620

615

610

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. 9

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Phe Met Asp Val Cys Met Pro Gly Val Glu Asn Tyr Gln Ile Ala Leu 660 665 670

Arg Ile His Glu Lys Phe Thr Lys Gln Arg His Gln Arg Pro Leu Leu 675 680 685

Val Ala Leu Ser Gly Asn Thr Asp Lys Ser Thr Lys Glu Lys Cys Met 690 695 700

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Tyr Gly Ala Thr His Leu Ile Asn Leu Trp Thr Phe Thr Thr His Ser 65 70 75 80

Arg Thr Val Ala Leu Val Met Thr Thr Ala Lys Val Leu Thr Ala Val 85 90 95

Val Ser Cys Ala Thr Ala Leu Met Leu Val His Ile Ile Pro Asp Leu
100 105 110

Leu Ser Val Lys Thr Arg Glu Leu Phe Leu Lys Asn Lys Ala Ala Glu 115 120 125

Leu Asp Arg Glu Met Gly Leu Ile Arg Thr Gln Glu Glu Thr Gly Arg 135 His Val Arg Met Leu Thr His Glu Ile Arg Ser Thr Leu Asp Arg His 150 155 Thr Ile Leu Lys Thr Thr Leu Val Glu Leu Gly Arg Thr Leu Ala Leu 165 170 Glu Glu Cys Ala Leu Trp Met Pro Thr Arg Thr Gly Leu Glu Leu Gln 185 Leu Ser Tyr Thr Leu Arg His Gln His Pro Val Glu Tyr Thr Val Pro 195 200 Ile Gln Leu Pro Val Ile Asn Gln Val Phe Gly Thr Ser Arg Ala Val 215 Lys Ile Ser Pro Asn Ser Pro Val Ala Arg Leu Arg Pro Val Ser Gly 230 235 Lys Tyr Met Leu Gly Glu Val Val Ala Val Arg Val Pro Leu Leu His 245 250 Leu Ser Asn Phe Gln Ile Asn Asp Trp Pro Glu Leu Ser Thr Lys Arg 265 Tyr Ala Leu Met Val Leu Met Leu Pro Ser Asp Ser Ala Arg Gln Trp 280 His Val His Glu Leu Glu Leu Val Glu Val Val Ala Asp Gln Val Ala 295 Val Ala Leu Ser His Ala Ala Ile Leu Glu Glu Ser Met Arg Ala Arg 310 315 Asp Leu Leu Met Glu Gln Asn Val Ala Leu Asp Leu Ala Arg Arg Glu 325 330 Ala Glu Thr Ala Ile Arg Ala Arg Asn Asp Phe Leu Ala Val Met Asn 345 His Glu Met Arg Thr Pro Met His Ala Ile Ile Ala Leu Ser Ser Leu 355 Leu Gln Glu Thr Glu Leu Thr Pro Glu Gln Arg Leu Met Val Glu Thr 375 Ile Leu Lys Ser Ser Asn Leu Leu Ala Thr Leu Met Asn Asp Val Leu 390 395 Asp Leu Ser Arg Leu Glu Asp Gly Ser Leu Gln Leu Glu Leu Gly Thr 405 410

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- Leu Pro Glu Phe Val Val Gly Asp Glu Lys Arg Leu Met Gln Ile Ile 450 455 460
- Leu Asn Ile Val Gly Asn Ala Val Lys Phe Ser. Lys Gln Gly Ser Ile 465 470 475 480
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- Phe Asp Val Lys Leu Gly Ile Ser Glu Arg Ser Asn Glu Ser Lys Gln 580 585 590
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- Gly Leu Lys Val Leu Val Met Asp Glu Asn Gly Val Ser Arg Met Val 610 615 620
- Thr Lys Gly Leu Leu Val His Leu Gly Cys Glu Val Thr Thr Val Ser 625 630 635 640
- Ser Asn Glu Glu Cys Leu Arg Val Val Ser His Glu His Lys Val Val 645 650 655
- Phe Met Asp Val Cys Met Pro Gly Val Glu Asn Tyr Gln Ile Ala Leu 660 665 670
- Arg Ile His Glu Lys Phe Thr Lys Gln Arg His Gln Arg Pro Leu Leu 675 680 685
- Val Ala Leu Ser Gly Asn Thr Asp Lys Ser Thr Lys Glu Lys Cys Met 690 695 700
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	_	tgt Cys	_						_						_	576
		tat Tyr 195			_								_	_		624
		tta Leu	_									_		_	_	672
		tct Ser						_		-	_		_			720
		atg Met					_	_			_	_				768
		aat Asn												_	_	816
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		cat His														912
		ctc Leu														960
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					agt Ser									1536
					aat Asn									1584
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	_				atc Ile		_	_		_			_	1776

-		ata Ile 595	_		_		_		_						182	24
		aag Lys													187	72
		gga Gly													192	20
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- Tyr Thr Thr Leu Val Glu Leu Ser Lys Thr Leu Gly Leu Gln Asn Cys
 195 200 205

Met Leu Thr Gln Glu Ile Arg Lys Ser Leu Asp Arg His Thr Ile Leu

- Ala Val Trp Met Pro Asn Asp Gly Gly Thr Glu Met Asp Leu Thr His 210 215 220
- Glu Leu Arg Gly Arg Gly Gly Tyr Gly Gly Cys Ser Val Ser Met Glu 225 230 235 240
- Asp Leu Asp Val Val Arg Ile Arg Glu Ser Asp Glu Val Asn Val Leu 245 250 255
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- Ser Glu Ile Gly Ala Val Ala Ala Ile Arg Met Pro Met Leu Arg Val 275 280 285
- Ser Asp Phe Asn Gly Glu Leu Ser Tyr Ala Ile Leu Val Cys Val Leu 290 295 300
- Pro Gly Gly Thr Pro Arg Asp Trp Thr Tyr Gln Glu Ile Glu Ile Val 305 310 315 320
- Lys Val Val Ala Asp Gln Val Thr Val Ala Leu Asp His Ala Ala Val 325 330 335

- Leu Glu Glu Ser Gln Leu Met Arg Glu Lys Leu Ala Glu Gln Asn Arg 340 345 350
- Ala Leu Gln Met Ala Lys Arg Asp Ala Leu Arg Ala Ser Gln Ala Arg 355 360 365
- Asn Ala Phe Gln Lys Thr Met Ser Glu Gly Met Arg Arg Pro Met His 370 375 380
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- Val Phe Gln Val Ile Leu His Ile Val Gly Ser Leu Val Lys Pro Arg 485 490 495
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- Gln Asp Leu Ser Phe Gly Val Cys Lys Lys Val Val Gln Leu Ile His 580 585 590
- Gly Asn Ile Ser Val Val Pro Gly Ser Asp Gly Ser Pro Glu Thr Met
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- Ser Leu Leu Arg Phe Arg Arg Pro Ser Ile Ser Val His Gly
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- Ser Ser Glu Ser Pro Ala Pro Asp His His Ala His Pro His Ser Asn

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Thr	Ala	Val 675	Ser	Ser	Gly	Phe	Asp 680	Cys	Leu	Thr	Ala	Ile 685	Ala	Pro	Gly	
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23 · UF-325XC1

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